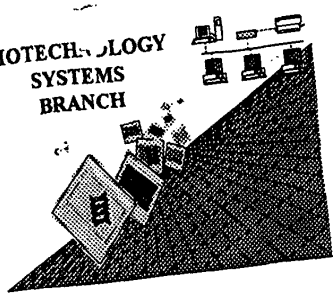
RAW SEQUENCE LISTING ERROR REPORT



The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 09/825,561

Source: OIPE

Date Processed by STIC: 5/30/2001

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.
PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:
1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY
FOR CRF SUBMISSION QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4216.
PATENTIN 2.1 e-mail help: patin21help@uspto.gov or phone 703-306-4119 (R. Wax)
PATENTIN 3.0 e-mail help: patin3help@uspto.gov or phone 703-306-4119 (R. Wax)

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 3.0 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW:

Checker Version 3.0

The Checker Version 3.0 application is a state-of the-art Windows based software program employing a logical and intuitive user-interface to check whether a sequence listing is in compliance with format and content rules. Checker Version 3.0 works for sequence listings generated for the original version of 37 CFR §§1.821 – 1.825 effective October 1, 1990 (old rules) and the revised version (new rules) effective July 1, 1998 as well as World Intellectual Property Organization (WIPO) Standard ST.25.
Checker Version 3.0 replaces the previous DOS-based version of Checker, and is Y2K-compliant. Checker allows public users to check sequence listings in Computer Readable form (CRF) before submitting them to the United States Patent and Trademark Office (USPTO). Use of Checker prior to filing the sequence listing is expected to result in fewer errored sequence listings, thus saving time and money.

Checker Version 3.0 can be down loaded from the USPTO website at the following address:
http://www.uspto.gov/web/offices/pac/checker

OIPE

RAW SEQUENCE LISTING DATE: 05/30/2001
PATENT APPLICATION: US/09/825,561 TIME: 15:13:21

Input Set : A:\00-22.SEQ.txt
Output Set: C:\CRF3\05302001\I825561.raw

Does Not Comply
Corrected Diskette Needed

```
     4 <110> APPLICANT: Sprecher, Cindy A.
     5       Novak, Julia E.
     6       West, James W.
     7       Presnell, Scott R.
     8       Holly, Richard D.
     9       Nelson, Andrew J.
    11 <120> TITLE OF INVENTION: SOLUBLE ZALPHA11 CYTOKINE RECEPTORS
    13 <130> FILE REFERENCE: 00-22
C--> 15 <140> CURRENT APPLICATION NUMBER: US/09/825,561
C--> 15 <141> CURRENT FILING DATE: 2001-04-03
    15 <150> PRIOR APPLICATION NUMBER: US 60/194,731
    16 <151> PRIOR FILING DATE: 2000-04-05
    18 <150> PRIOR APPLICATION NUMBER: US 60/222,121
    19 <151> PRIOR FILING DATE: 2000-07-28
    21 <160> NUMBER OF SEQ ID NOS: 86
    23 <170> SOFTWARE: FastSEQ for Windows Version 3.0
    25 <210> SEQ ID NO: 1
    26 <211> LENGTH: 1614
    27 <212> TYPE: DNA
    28 <213> ORGANISM: Homo sapiens
    30 <220> FEATURE:
    31 <221> NAME/KEY: CDS
    32 <222> LOCATION: (1)...(1614)
    34 <400> SEQUENCE: 1
    35  atg ccg cgt ggc tgg gcc gcc ccc ttg ctc ctg ctg ctg ctc cag gga       48
    36  Met Pro Arg Gly Trp Ala Ala Pro Leu Leu Leu Leu Leu Gln Gly
    37   1               5                  10                  15
    39  ggc tgg ggc tgc ccc gac ctc gtc tgc tac acc gat tac ctc cag acg       96
    40  Gly Trp Gly Cys Pro Asp Leu Val Cys Tyr Thr Asp Tyr Leu Gln Thr
    41              20                  25                  30
    43  gtc atc tgc atc ctg gaa atg tgg aac ctc cac ccc agc acg ctc acc      144
    44  Val Ile Cys Ile Leu Glu Met Trp Asn Leu His Pro Ser Thr Leu Thr
    45          35                  40                  45
    47  ctt acc tgg caa gac cag tat gaa gag ctg aag gac gag gcc acc tcc      192
    48  Leu Thr Trp Gln Asp Gln Tyr Glu Glu Leu Lys Asp Glu Ala Thr Ser
    49      50                  55                  60
    51  tgc agc ctc cac agg tcg gcc cac aat gcc acg cat gcc acc tac acc      240
    52  Cys Ser Leu His Arg Ser Ala His Asn Ala Thr His Ala Thr Tyr Thr
    53   65                  70                  75                  80
    55  tgc cac atg gat gta ttc cac ttc atg gcc gac gac att ttc agt gtc      288
    56  Cys His Met Asp Val Phe His Phe Met Ala Asp Asp Ile Phe Ser Val
    57                  85                  90                  95
    59  aac atc aca gac cag tct ggc aac tac tcc cag gag tgt ggc agc ttt      336
    60  Asn Ile Thr Asp Gln Ser Gly Asn Tyr Ser Gln Glu Cys Gly Ser Phe
    61              100                 105                 110
    63  ctc ctg gct gag agc atc aag ccg gct ccc cct ttc aac gtg act gtg      384
    64  Leu Leu Ala Glu Ser Ile Lys Pro Ala Pro Pro Phe Asn Val Thr Val
```

RAW SEQUENCE LISTING DATE: 05/30/2001
PATENT APPLICATION: **US/09/825,561** TIME: 15:13:21

Input Set : **A:\00-22.SEQ.txt**
Output Set: **C:\CRF3\05302001\I825561.raw**

```
65          115                 120                 125
67  acc ttc tca gga cag tat aat atc tcc tgg cgc tca gat tac gaa gac      432
68  Thr Phe Ser Gly Gln Tyr Asn Ile Ser Trp Arg Ser Asp Tyr Glu Asp
69      130                 135                 140
71  cct gcc ttc tac atg ctg aag ggc aag ctt cag tat gag ctg cag tac      480
72  Pro Ala Phe Tyr Met Leu Lys Gly Lys Leu Gln Tyr Glu Leu Gln Tyr
73  145                 150                 155                 160
75  agg aac cgg gga gac ccc tgg gct gtg agt ccg agg aga aag ctg atc      528
76  Arg Asn Arg Gly Asp Pro Trp Ala Val Ser Pro Arg Arg Lys Leu Ile
77                  165                 170                 175
79  tca gtg gac tca aga agt gtc tcc ctc ctc ccc ctg gag ttc cgc aaa      576
80  Ser Val Asp Ser Arg Ser Val Ser Leu Leu Pro Leu Glu Phe Arg Lys
81              180                 185                 190
83  gac tcg agc tat gag ctg cag gtg cgg gca ggg ccc atg cct ggc tcc      624
84  Asp Ser Ser Tyr Glu Leu Gln Val Arg Ala Gly Pro Met Pro Gly Ser
85          195                 200                 205
87  tcc tac cag ggg acc tgg agt gaa tgg agt gac ccg gtc atc ttt cag      672
88  Ser Tyr Gln Gly Thr Trp Ser Glu Trp Ser Asp Pro Val Ile Phe Gln
89      210                 215                 220
91  acc cag tca gag gag tta aag gaa ggc tgg aac cct cac ctg ctg ctt      720
92  Thr Gln Ser Glu Glu Leu Lys Glu Gly Trp Asn Pro His Leu Leu Leu
93  225                 230                 235                 240
95  ctc ctc ctg ctt gtc ata gtc ttc att cct gcc ttc tgg agc ctg aag      768
96  Leu Leu Leu Leu Val Ile Val Phe Ile Pro Ala Phe Trp Ser Leu Lys
97                  245                 250                 255
99  acc cat cca ttg tgg agg cta tgg aag aag ata tgg gcc gtc ccc agc      816
100  Thr His Pro Leu Trp Arg Leu Trp Lys Lys Ile Trp Ala Val Pro Ser
101             260                 265                 270
103  cct gag cgg ttc ttc atg ccc ctg tac aag ggc tgc agc gga gac ttc      864
104  Pro Glu Arg Phe Phe Met Pro Leu Tyr Lys Gly Cys Ser Gly Asp Phe
105         275                 280                 285
107  aag aaa tgg gtg ggt gca ccc ttc act ggc tcc agc ctg gag ctg gga      912
108  Lys Lys Trp Val Gly Ala Pro Phe Thr Gly Ser Ser Leu Glu Leu Gly
109      290                 295                 300
111  ccc tgg agc cca gag gtg ccc tcc acc ctg gag gtg tac agc tgc cac      960
112  Pro Trp Ser Pro Glu Val Pro Ser Thr Leu Glu Val Tyr Ser Cys His
113  305                 310                 315                 320
115  cca cca cgg agc ccg gcc aag agg ctg cag ctc acg gag cta caa gaa     1008
116  Pro Pro Arg Ser Pro Ala Lys Arg Leu Gln Leu Thr Glu Leu Gln Glu
117                 325                 330                 335
119  cca gca gag ctg gtg gag tct gac ggt gtg ccc aag ccc agc ttc tgg     1056
120  Pro Ala Glu Leu Val Glu Ser Asp Gly Val Pro Lys Pro Ser Phe Trp
121             340                 345                 350
123  ccg aca gcc cag aac tcg ggg ggc tca gct tac agt gag gag agg gat     1104
124  Pro Thr Ala Gln Asn Ser Gly Gly Ser Ala Tyr Ser Glu Glu Arg Asp
125         355                 360                 365
127  cgg cca tac ggc ctg gtg tcc att gac aca gtg act gtg cta gat gca     1152
128  Arg Pro Tyr Gly Leu Val Ser Ile Asp Thr Val Thr Val Leu Asp Ala
129      370                 375                 380
```

RAW SEQUENCE LISTING DATE: 05/30/2001
PATENT APPLICATION: **US/09/825,561** TIME: 15:13:21

Input Set : **A:\00-22.SEQ.txt**
Output Set: **C:\CRF3\05302001\I825561.raw**

```
131  gag ggg cca tgc acc tgg ccc tgc agc tgt gag gat gac ggc tac cca     1200
132  Glu Gly Pro Cys Thr Trp Pro Cys Ser Cys Glu Asp Asp Gly Tyr Pro
133  385                 390                 395                 400
135  gcc ctg gac ctg gat gct ggc ctg gag ccc agc cca ggc cta gag gac     1248
136  Ala Leu Asp Leu Asp Ala Gly Leu Glu Pro Ser Pro Gly Leu Glu Asp
137                  405                 410                 415
139  cca ctc ttg gat gca ggg acc aca gtc ctg tcc tgt ggc tgt gtc tca     1296
140  Pro Leu Leu Asp Ala Gly Thr Thr Val Leu Ser Cys Gly Cys Val Ser
141              420                 425                 430
143  gct ggc agc cct ggg cta gga ggg ccc ctg gga agc ctc ctg gac aga     1344
144  Ala Gly Ser Pro Gly Leu Gly Gly Pro Leu Gly Ser Leu Leu Asp Arg
145          435                 440                 445
147  cta aag cca ccc ctt gca gat ggg gag gac tgg gct ggg gga ctg ccc     1392
148  Leu Lys Pro Pro Leu Ala Asp Gly Glu Asp Trp Ala Gly Gly Leu Pro
149      450                 455                 460
151  tgg ggt ggc cgg tca cct gga ggg gtc tca gag agt gag gcg ggc tca     1440
152  Trp Gly Gly Arg Ser Pro Gly Gly Val Ser Glu Ser Glu Ala Gly Ser
153  465                 470                 475                 480
155  ccc ctg gcc ggc ctg gat atg gac acg ttt gac agt ggc ttt gtg ggc     1488
156  Pro Leu Ala Gly Leu Asp Met Asp Thr Phe Asp Ser Gly Phe Val Gly
157                  485                 490                 495
159  tct gac tgc agc agc cct gtg gag tgt gac ttc acc agc ccc ggg gac     1536
160  Ser Asp Cys Ser Ser Pro Val Glu Cys Asp Phe Thr Ser Pro Gly Asp
161              500                 505                 510
163  gaa gga ccc ccc cgg agc tac ctc cgc cag tgg gtg gtc att cct ccg     1584
164  Glu Gly Pro Pro Arg Ser Tyr Leu Arg Gln Trp Val Val Ile Pro Pro
165          515                 520                 525
167  cca ctt tcg agc cct gga ccc cag gcc agc                           1614
168  Pro Leu Ser Ser Pro Gly Pro Gln Ala Ser
169      530                 535
171 <210> SEQ ID NO: 2
172 <211> LENGTH: 538
173 <212> TYPE: PRT
174 <213> ORGANISM: Homo sapiens
176 <400> SEQUENCE: 2
177  Met Pro Arg Gly Trp Ala Ala Pro Leu Leu Leu Leu Leu Leu Gln Gly
178   1               5                  10                  15
179  Gly Trp Gly Cys Pro Asp Leu Val Cys Tyr Thr Asp Tyr Leu Gln Thr
180              20                  25                  30
181  Val Ile Cys Ile Leu Glu Met Trp Asn Leu His Pro Ser Thr Leu Thr
182          35                  40                  45
183  Leu Thr Trp Gln Asp Gln Tyr Glu Glu Leu Lys Asp Glu Ala Thr Ser
184      50                  55                  60
185  Cys Ser Leu His Arg Ser Ala His Asn Ala Thr His Ala Thr Tyr Thr
186  65                  70                  75                  80
187  Cys His Met Asp Val Phe His Phe Met Ala Asp Asp Ile Phe Ser Val
188                  85                  90                  95
189  Asn Ile Thr Asp Gln Ser Gly Asn Tyr Ser Gln Glu Cys Gly Ser Phe
190              100                 105                 110
```

RAW SEQUENCE LISTING DATE: 05/30/2001
PATENT APPLICATION: **US/09/825,561** TIME: 15:13:21

Input Set : **A:\00-22.SEQ.txt**
Output Set: **C:\CRF3\05302001\I825561.raw**

```
191  Leu Leu Ala Glu Ser Ile Lys Pro Ala Pro Pro Phe Asn Val Thr Val
192          115                 120                 125
193  Thr Phe Ser Gly Gln Tyr Asn Ile Ser Trp Arg Ser Asp Tyr Glu Asp
194      130                 135                 140
195  Pro Ala Phe Tyr Met Leu Lys Gly Lys Leu Gln Tyr Glu Leu Gln Tyr
196  145                 150                 155                 160
197  Arg Asn Arg Gly Asp Pro Trp Ala Val Ser Pro Arg Arg Lys Leu Ile
198                  165                 170                 175
199  Ser Val Asp Ser Arg Ser Val Ser Leu Leu Pro Leu Glu Phe Arg Lys
200              180                 185                 190
201  Asp Ser Ser Tyr Glu Leu Gln Val Arg Ala Gly Pro Met Pro Gly Ser
202          195                 200                 205
203  Ser Tyr Gln Gly Thr Trp Ser Glu Trp Ser Asp Pro Val Ile Phe Gln
204      210                 215                 220
205  Thr Gln Ser Glu Glu Leu Lys Glu Gly Trp Asn Pro His Leu Leu Leu
206  225                 230                 235                 240
207  Leu Leu Leu Leu Val Ile Val Phe Ile Pro Ala Phe Trp Ser Leu Lys
208                  245                 250                 255
209  Thr His Pro Leu Trp Arg Leu Trp Lys Lys Ile Trp Ala Val Pro Ser
210              260                 265                 270
211  Pro Glu Arg Phe Phe Met Pro Leu Tyr Lys Gly Cys Ser Gly Asp Phe
212          275                 280                 285
213  Lys Lys Trp Val Gly Ala Pro Phe Thr Gly Ser Ser Leu Glu Leu Gly
214      290                 295                 300
217  Pro Trp Ser Pro Glu Val Pro Ser Thr Leu Glu Val Tyr Ser Cys His
218  305                 310                 315                 320
219  Pro Pro Arg Ser Pro Ala Lys Arg Leu Gln Leu Thr Glu Leu Gln Glu
220                  325                 330                 335
221  Pro Ala Glu Leu Val Glu Ser Asp Gly Val Pro Lys Pro Ser Phe Trp
222              340                 345                 350
223  Pro Thr Ala Gln Asn Ser Gly Gly Ser Ala Tyr Ser Glu Glu Arg Asp
224          355                 360                 365
225  Arg Pro Tyr Gly Leu Val Ser Ile Asp Thr Val Thr Val Leu Asp Ala
226      370                 375                 380
227  Glu Gly Pro Cys Thr Trp Pro Cys Ser Cys Glu Asp Asp Gly Tyr Pro
228  385                 390                 395                 400
229  Ala Leu Asp Leu Asp Ala Gly Leu Glu Pro Ser Pro Gly Leu Glu Asp
230                  405                 410                 415
231  Pro Leu Leu Asp Ala Gly Thr Thr Val Leu Ser Cys Gly Cys Val Ser
232              420                 425                 430
233  Ala Gly Ser Pro Gly Leu Gly Gly Pro Leu Gly Ser Leu Leu Asp Arg
234          435                 440                 445
235  Leu Lys Pro Pro Leu Ala Asp Gly Glu Asp Trp Ala Gly Gly Leu Pro
236      450                 455                 460
237  Trp Gly Gly Arg Ser Pro Gly Gly Val Ser Glu Ser Glu Ala Gly Ser
238  465                 470                 475                 480
239  Pro Leu Ala Gly Leu Asp Met Asp Thr Phe Asp Ser Gly Phe Val Gly
240                  485                 490                 495
241  Ser Asp Cys Ser Ser Pro Val Glu Cys Asp Phe Thr Ser Pro Gly Asp
```

RAW SEQUENCE LISTING DATE: 05/30/2001
PATENT APPLICATION: **US/09/825,561** TIME: 15:13:21

Input Set : **A:\00-22.SEQ.txt**
Output Set: **C:\CRF3\05302001\I825561.raw**

```
242              500                 505                 510
243  Glu Gly Pro Pro Arg Ser Tyr Leu Arg Gln Trp Val Val Ile Pro Pro
244          515                 520                 525
245  Pro Leu Ser Ser Pro Gly Pro Gln Ala Ser
246      530                 535
248 <210> SEQ ID NO: 3
249 <211> LENGTH: 696
250 <212> TYPE: DNA
251 <213> ORGANISM: Homo sapiens
253 <220> FEATURE:
254 <221> NAME/KEY: CDS
255 <222> LOCATION: (1)...(696)
257 <400> SEQUENCE: 3
258  ctg aac acg aca att ctg acg ccc aat ggg aat gaa gac acc aca gct       48
259  Leu Asn Thr Thr Ile Leu Thr Pro Asn Gly Asn Glu Asp Thr Thr Ala
260   1               5                  10                  15
262  gat ttc ttc ctg acc act atg ccc act gac tcc ctc agt gtt tcc act       96
263  Asp Phe Phe Leu Thr Thr Met Pro Thr Asp Ser Leu Ser Val Ser Thr
264              20                  25                  30
266  ctg ccc ctc cca gag gtt cag tgt ttt gtg ttc aat gtc gag tac atg      144
267  Leu Pro Leu Pro Glu Val Gln Cys Phe Val Phe Asn Val Glu Tyr Met
268          35                  40                  45
270  aat tgc act tgg aac agc agc tct gag ccc cag cct acc aac ctc act      192
271  Asn Cys Thr Trp Asn Ser Ser Ser Glu Pro Gln Pro Thr Asn Leu Thr
272       50                  55                  60
274  ctg cat tat tgg tac aag aac tcg gat aat gat aaa gtc cag aag tgc      240
275  Leu His Tyr Trp Tyr Lys Asn Ser Asp Asn Asp Lys Val Gln Lys Cys
276   65                  70                  75                  80
278  agc cac tat cta ttc tct gaa gaa atc act tct ggc tgt cag ttg caa      288
279  Ser His Tyr Leu Phe Ser Glu Glu Ile Thr Ser Gly Cys Gln Leu Gln
280                  85                  90                  95
282  aaa aag gag atc cac ctc tac caa aca ttt gtt gtt cag ctc cag gac      336
283  Lys Lys Glu Ile His Leu Tyr Gln Thr Phe Val Val Gln Leu Gln Asp
284              100                 105                 110
286  cca cgg gaa ccc agg aga cag gcc aca cag atg cta aaa ctg cag aat      384
287  Pro Arg Glu Pro Arg Arg Gln Ala Thr Gln Met Leu Lys Leu Gln Asn
288          115                 120                 125
290  ctg gtg atc ccc tgg gct cca gag aac cta aca ctt cac aaa ctg agt      432
291  Leu Val Ile Pro Trp Ala Pro Glu Asn Leu Thr Leu His Lys Leu Ser
292      130                 135                 140
294  gaa tcc cag cta gaa ctg aac tgg aac aac aga ttc ttg aac cac tgt      480
295  Glu Ser Gln Leu Glu Leu Asn Trp Asn Asn Arg Phe Leu Asn His Cys
296  145                 150                 155                 160
298  ttg gag cac ttg gtg cag tac cgg act gac tgg gac cac agc tgg act      528
299  Leu Glu His Leu Val Gln Tyr Arg Thr Asp Trp Asp His Ser Trp Thr
300                  165                 170                 175
302  gaa caa tca gtg gat tat aga cat aag ttc tcc ttg cct agt gtg gat      576
303  Glu Gln Ser Val Asp Tyr Arg His Lys Phe Ser Leu Pro Ser Val Asp
304              180                 185                 190
```

<210> 16
<211> 567
<212> PRT
<213> Artificial Sequence

see item 11 on Error Summary Sheet

<400> 16
Met Pro Arg Gly Trp Ala Ala Pro Leu Leu Leu Leu Leu Leu Gln Gly
1 5 10 15

The types of errors shown exist <u>throughout</u> the Sequence Listing. Please check subsequent sequences for similar errors.

FYI

<u>Please Note:</u>
Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

VERIFICATION SUMMARY DATE: 05/30/2001
PATENT APPLICATION: **US/09/825,561** TIME: 15:13:22

Input Set : **A:\00-22.SEQ.txt**
Output Set: **C:\CRF3\05302001\I825561.raw**

L:15 M:270 C: Current Application Number differs, Replaced Current Application No
L:15 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:476 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7
L:477 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7
L:478 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7
L:479 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7
L:480 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7
L:481 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7
L:482 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7
L:483 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7
L:484 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7
L:485 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7
L:486 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7
L:502 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8
L:503 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8
L:504 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8
L:505 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8
L:506 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8
L:507 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8
L:508 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8
L:509 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8
L:510 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8
L:511 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8
L:512 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8
L:513 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8
L:838 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:13
L:1018 M:258 W: Mandatory Feature missing, <220> FEATURE:
L:1018 M:258 W: Mandatory Feature missing, <223> OTHER INFORMATION:
L:1207 M:258 W: Mandatory Feature missing, <220> FEATURE:
L:1207 M:258 W: Mandatory Feature missing, <223> OTHER INFORMATION:
L:1560 M:258 W: Mandatory Feature missing, <220> FEATURE:
L:1560 M:258 W: Mandatory Feature missing, <223> OTHER INFORMATION:
L:1645 M:258 W: Mandatory Feature missing, <220> FEATURE:
L:1645 M:258 W: Mandatory Feature missing, <223> OTHER INFORMATION:
L:2365 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:66
L:2366 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:66
L:2367 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:66
L:2368 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:66
L:2369 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:66
L:2370 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:66
L:2371 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:66
L:2372 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:66
L:2373 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:66
L:2374 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:66
L:2375 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:66
L:2376 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:66
L:2377 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:66
L:2392 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:67

VERIFICATION SUMMARY DATE: 05/30/2001
PATENT APPLICATION: **US/09/825,561** TIME: 15:13:22

Input Set : **A:\00-22.SEQ.txt**
Output Set: **C:\CRF3\05302001\I825561.raw**

L:2554 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:71
L:2556 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:71
L:2720 M:258 W: Mandatory Feature missing, <220> FEATURE:
L:2720 M:258 W: Mandatory Feature missing, <223> OTHER INFORMATION: